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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/760,320	01/21/2004	Takao Isogai	084335-0181	7661
	7590 01/30/200 LARDNER LLP	EXAMINER MARTINELL, JAMES		
SUITE 500	T NIW			
3000 K STREE WASHINGTO			ART UNIT	PAPER NUMBER
			1634	
SHORTENED STATUTOR	Y PERIOD OF RESPONSE	MAIL DATE	DELIVERY MODE	
3 MOI	NTHS	01/30/2007	PAPER	

Please find below and/or attached an Office communication concerning this application or proceeding.

If NO period for reply is specified above, the maximum statutory period will apply and will expire 6 MONTHS from the mailing date of this communication.

U.S. Patent and Trademark Office PTOL-326 (Rev. 08-06)

1) Notice of References Cited (PTO-892)

Paper No(s)/Mail Date _

2) Notice of Draftsperson's Patent Drawing Review (PTO-948)

3) Information Disclosure Statement(s) (PTO/SB/08)

4) Interview Summary (PTO-413)
Paper No(s)/Mail Date. _____.

6) Other:

Application/Control Number: 10/760,320

Art Unit: 1634

The text of those sections of Title 35, U.S. Code not included in this action can be found in a prior Office action.

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Claims 1 and 5-7 are are rejected under 35 U.S.C. 102(b) as being clearly anticipated by GenBank® Accession No. AC008736 (September 27, 2000). GenBank® Accession No. AC008736 has 92.7% sequence identity to SEQ ID NO: 102 (see the alignment below). Thus, GenBank® Accession No. AC008736 is embraced by the claims (*e.g.*, see Claim 1(f) and (g)). Since the DNA was sequenced, it was necessarily contained within a vector and host cell. This rejection is repeated for reasons already of record (*e.g.*, Office action mailed June 2, 2006, pages 5-10). Applicants' arguments (response filed October 25, 2006, pages 10-11) are most unconvincing. SEQ ID NO: 2290 is encoded by nucleotides 858-1562 of SEQ ID NO: 102 (see the alignment below). GenBank® Accession No. AC008736 matches SEQ ID NO: 102, nucleotides 858-1562 except for one mismatch at nucleotide 1187 in SEQ ID NO: 102 (see the alignment below). SEQ ID NO: 102, nucleotides 1185-1187 (cga) encode arginine as do GenBank® Accession No. AC008736, nucleotides 128,245-128,247 (cgg). Thus, the nucleic acid of GenBank® Accession No. AC008736 is embraced by the claims. Watson et al (*The DNA Story*, 1981 W.H. Freeman and Company, New York, page 547) is cited here to show that both cga and cgg encode arginine.

RESULT 3 AC008736/c

LOCUS AC008736 191925 bp DNA linear PRI 27-SEP-2000

DEFINITION Homo sapiens chromosome 19 clone CTD-2538C1, complete sequence.

ACCESSION AC008736

VERSION AC008736.6 GI:10312244

KEYWORDS HTG.

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;

Hominidae; Homo.

REFERENCE 1 (bases 1 to 191925)

AUTHORS DOE Joint Genome Institute and Stanford Human Genome Center.

TITLE Direct Submission

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 191925)
AUTHORS DOE Joint Genome Institute.

TITLE Direct Submission

JOURNAL Submitted (03-AUG-1999) Production Sequencing Facility, DOE Joint

```
Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA
REFERENCE
          (bases 1 to 191925)
 AUTHORS
        DOE Joint Genome Institute and Stanford Human Genome Center.
 TITLE
        Direct Submission
 JOURNAL
        Submitted (27-SEP-2000) DOE Joint Genome Institute, 2800 Mitchell
        Drive, Walnut Creek, CA 94598, USA
COMMENT
        On Sep 27, 2000 this sequence version replaced gi:8575905.
        Draft Sequence Produced by DOE Joint Genome Institute
        www.jqi.doe.gov
        Finishing Completed at Stanford Human Genome Center.
        www-shqc.stanford.edu
        Quality: Phrap Quality >=40 99.9% of Sequence;
        Estimated Total Number of Errors is 0.1.
        STS Content:
        SHGC-57769 G37408.
FEATURES
               Location/Qualifiers
               1. .191925
   source
               /organism="Homo sapiens"
               /mol type="genomic DNA"
               /db_xref="taxon:9606"
               /chromosome="19"
               /clone="CTD-2538C1"
ORIGIN
 Query Match
                  92.7%; Score 2892.8; DB 8; Length 191925;
 Best Local Similarity
                  99:9%;
                        Pred. No. 0;
 Matches 2894; Conservative
                       0; Mismatches
                                       Indels
                                              0;
                                                 Gaps
                                                       0:
Qу
        1 ACTAGAGGTGGGGTTAGCGCTTGGAAGCACCGACCAACGTGAGCGCAACGCGGCAGGGAC 60
          129460 ACTAGAGGTGGGGTTAGCGCTTGGAAGCACCGACCAACGTGAGCGCAACGCGGCAGGGAC 129401
Db
Qу
        61 ACCTGACCCCGGCGGCGCCCAGCCCCTCGGATTGCCAGTCACTGCTCGCTTTGGGGCACG 120
          129400 ACCTGACCCCGGCGCGCCCCAGCCCCTCGGATTGCCAGTCACTGCTCGCTTTGGGGCACG 129341
Db
       121 GAGGTGCCCAGTCCTGCGGGGCACCCGACGTCCTGTCGCCGACAGGGTCCGGGAGTCAGT 180
Qу
          129340 GAGGTGCCCAGTCCTGCGGGGCACCCGACGTCCTGTCGCCGACAGGGTCCGGGAGTCAGT 129281
Db
       181 ATAGCTGGGTTCTAGTCCCATCACAGGCAAAAACTCCGCGGGAGCCTGGCCCGCTTTTTA 240
Qу
          129280 ATAGCTGGGTTCTAGTCCCATCACAGGCAAAAACTCCGCGGGAGCCTGGCCCGCTTTTTA 129221
Db
       241 CCTGGGCCTCAGTTTCCCCATCCGTAAAATAGAACGGGTTGGATCTCCCGAGCGCTAACA 300
Qу
          Db
     129220 CCTGGGCCTCAGTTTCCCCATCCGTAAAATAGAACGGGTTGGATCTCCCGAGCGCTAACA 129161
       Qу
          Db
       Qу
          Db
       421 GTGGGCGGTCCTAGGAAACCCTACCGGCCGCCCTTGGCAGCGCCTAAGGCGGAĢCGCG 480
Qу
          129040 GTGGGCGGTCCTAGGAAACCCTACCGGCCGCCCTTGGCAGCGCCTAAGGCGGAGCGCG 128981
Db
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Qу	481	CGGCTCTGCAGCCTGCTTGCCCCGGAGTTGGCACCCACGGAGGATGGGGACCGCACCCTC	540
Db	128980	CGGCTCTGCAGCCTGCTTGCCCCGGAGTTGGCACCCACGGAGGATGGGGACCGCACCCTC	128921
Qу	541	AGCTTCGCAGGGAGCCACCGTGGAGGCCAGGGCGGTGCAGAGACACGACGTGTGACTCGG	600
Db	128920	AGCTTCGCAGGGAGCCACCGTGGAGGCCAGGGCGGTGCAGAGACACGACGTGTGACTCGG	128861
Qу	601	AGTGCGCCTGGGGAGGATGGACGAGGGAGCGGGGGACCGCTAACGGGGCTCCCTCTGCGC	660
Db	128860	AGTGCGCCTGGGGAGGATGGACGAGGGAGCGGGGACCGCTAACGGGGCTCCCTCTGCGC	128801
Qу	661	GCCCGTCCGCAGAGGCGCACGTCGAGGGTCCCGGGCGGCTCCGTGGACGTTGGCGGTA	720
Db	128800	GCCCGTCCGCAGAGGCCACGTCGAGGGTCCCGGGCGGCTCCGTGGACGTTGGCGGTA	128741
Qу	. 721	GCGCCGAGCGACCATGAAGAGCGTTCGTGCCGCGCGCCCAAGGCCGGGATG	780
Db	128740	GCGCCGAGCCATGAAGAGCGTTCGTGCCGCGCGCCCAAGGCCGGGATG	128681
Qy		GGGGTTAGCCACATCCTGCCGCGCTGAGGGGGGGGGCCTAACGGGCGGG	840
Db		GGGGTTAGCCACATCCTGCCGCGCTGAGGGGGAGGCTAACGGGCGCGGGCCGGCC	128621
Qу	841	AGCCGGAGCCCACCGCGATGGCGAGGGAGGAGTGCAAGGCGCTGCTGGACGGGCTCAACA	900
Db	128620	AGCCGGAGCCCACCGCGATGCCGAGGGAGGAGTGCAAGGCGCTGCTGGACGGGCTCAACA	128561
Qу	901	AGACGACTGCGTGCTACCACCACCTGGTGCTGACCGTCGGTGGCTCGGCGGACTCGCAGA	960
Db	128560	AGACGACTGCGTGCTACCACCACCTGGTGCTGACCGTCGGTGGCTCGGCGGACTCGCAGA	128501
Qу	961	ACCTGCGGCAGGAGCTGCAAAAGACGCCCCAGAAGGCGCAGGAGCTGGCGGTGTCCACCT	1020
Db	128500	ACCTGCGGCAGGAGCTGCAAAAGACGCCCCAGAAGGCGCAGGAGCTGGCGGTGTCCACCT	128441
Qу	1021	GCGCCCGGCTGACTGCTGCTGCCGCGACCGGGGCCTGGCCGACGAGCGCGCCGAGT	1080
. Db	128440	GCGCCCGGCTGACTGCTGCTGCGCGACCGGGGCCTGGCCGACGAGCGCGCCGAGT	128381
QУ	1081	TCGAGCGGCTCTGGGTGGCCTTCTCGGGCTGCCTGGACCTGCTGGAAGCGGACATGCGAC	1140
Db	128380	${\tt TCGAGCGGCTCTGGGTGGCCTTCTCGGGCTGCCTGGACCTGCTGGAAGCGGACATGCGAC}$	128321
Qу	1141	GCTCGCTGGAGCTGGGCGCCGCGTTCCCGCTGCACGCGCGGCGACCGCTGGTGCGCA	1200
Db	128320	GCTCGCTGGAGCTGGGCGCCGCGTTCCCGCTGCACGCCGCGGCGGCCGCTGGTGCGCA	128261
ДУ	1201	CAGGTGTGGCGCGCCTCCTCCGGCGTGGCGCGCGCGCGCG	1260
Db	128260	CAGGTGTGGCGCCTCCTCCGGCGTGGCGCGCGCGCGCGCG	128201
Qу	1261	GGCTCGAGGCGGAGGGGGACTTCGACGTCGCGGACCTGCGGGAGCTGGAGCGCGAGGTCC	1320
Db	128200	GGCTCGAGGCGGAGCTTCGACGTCGCGGACCTGCGGAGCTGGAGCGCGAGGTCC	
Qу	1321	TTCAGGTGGGCGAGATGATCGACAACATGGAGATGAAGGTCAACGTGCCCCGCTGGACCG	1380
Db	128140	TTCAGGTGGCCGAGATGATCGACAACATGGAGATGAAGGTCAACGTGCCCCGCTGGACCG	128081

Qу	1381	$\tt TGCAAGCCCGGCAGGCGGGGGGGGGGGGGGGGGGGGGGG$	1440
Db	128080	TGCAAGCCCGGCAGGCGGCGGCCCGAGCTCCTGTCCACGGTCAGCGCCGGCCCCTCCT	128021
Qу	1441	CGGTCGTGTCCTTGCAGGAGCCCGGGGGGGGGGTTGCGACCCCAGGAAGGCCCTGGCCGCCA	1500
Db	128020	${\tt CGGTCGTGTCCTTGCAGGAGCGCGGGGGGGGGGTTGCGACCCCAGGAAGGCCCTGGCCGCCA}$	127961
Qу	1501	TCCTTTTCGGCGCCGTGCTGCCGGCGGCTGTGCCGTGGCGAAGCTGA	1560
Db	127960	TCCTTTTCGGCGCCGTGCTGCTGGCGGCGTGTGCCGTGTGCGTGGCGAAGCTGA	127901
Qу	1561	GCTGACAGACACCCGACGCCGCCTGCTGCTGCCGCTCCCTCGAGAAAAGACTCGG	1620
Db	127900	GCTGACAGACACCCGACGGCCGCCTGCTGCTGCCGCTCCCCTGAGAAAAGACTCGG	127841
Qу	1621	GATGGGTGTGGGGTCTGGCCTGTGCAAGGGGAGTGGTCCTAAAACCCCGTGTGTGCATGG	1680
Db	127840	GATGGGTGTGGGCCTGTGCAAGGGGAGTGGTCCTAAAACCCCGTGTGTGCATGG	127781
Qу	1681	GTACACGCGCTTTCCAGTGCACATCTGCCTGGGCAGGACACGGTTTTCCTCTTGCTGGC	1740
Db	127780	GTACACGCGCGTTTCCAGTGCACATCTGCCTGGGCAGGACACGGTTTTCCTCTTGCTGGC	127721
Qу	1741	CCGGGAGAAGTTAACTTTGCGCCGGCCGTCAGGGCATTACCGCTAACGTCTGCAGGAGCT	1800
Db	127720	CCGGGAGAAGTTAACTTTGCGCCGGCCGTCAGGGCATTACCGCTAACGTCTGCAGGAGCT	127661
Qу	1801	$\tt TTATTCCCTATTAATAGAAAACCGTCACAGTGACCCTAGATCCCTCCGAGTTAATGAGTT$	1860
Db	127660	TTATTCCCTATTAATAGAAAACCGTCACAGTGACCCTAGATCCCTCCGAGTTAATGAGTT	127601
Qу	1861	AACACATGTGCTGTTGGGGCGTCTTTACAGGGAGTCCGAGTTCGGTGCCCACCCCTGCCA	1920
Db	127600	AACACATGTGCTGTTGGGGCGTCTTTACAGGGAGTCCGAGTTCGGTGCCCACCCCTGCCA	127541
Qу	1921	GCGTCGCCCCTTTCTGCGTGGGACAGTTTGAAAAGGTGGGTG	1980
Db	127540	GCGTCGCCCCTTTCTGCGTGGGACAGTTTGAAAAGGTGGGTG	127481
Qy	1981	${\tt GAGAGGGACGCTGTTTGGTTCTATGTGGTTGGTTTTCCCGGACAAGAAAATTGCAA}$	2040
Db	127480	GAGAGGACGCTGTTTGGTTCTATGTGGTTGGTCTGTTTCCCGGACAAAAAATTGCAA	127421
Qу	2041	${\tt TCAAATGTCAGCAGCTTTTATTACCTTAATCTTTCAGGGCCTAAATTTAGGAGAGTGTCC}$	2100
Db	127420	TCAAATGTCAGCAGCTTTTATTACCTTAATCTTTCAGGGCCTAAATTTAGGAGAGTGTCC	127361
Qу	2101	${\tt TGAGAGCAGTTCATACAAAGGGCTTTCTCTAAGACGCGCTACAGCCCTTCCTAGCAGAGT}$	2160
Db	127360	TGAGAGCAGTTCATACAAAGGGCTTTCTCTAAGACGCGCTACAGCCCTTCCTAGCAGAGT	127301
Qу	2161	${\tt TTATCCATTCGTCCCCAAGAGCAGCTAGAAGAGATTTGAGGTCATGACCTCCCACTGCCG}$	2220
Db	127300		127241
Qy	2221	$\tt CTCAGGGGCTGACCCTATTTAGGAAACCAAAGAGGGTGGGT$	2280
Db	127240		127181

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Qу
    2281 TTGGATCCAGTGCGCACACTTGCCTGCGGAAAAGGGCTCTCCCCAGCCACCCGGAGATGG 2340
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Db
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       127120 GGGTAAGAGGAAGAGCAGAGGCTTGGGGTAGGGCCACCTGGTGTTTAAACAGGCACTTTC 127061
Db
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Qy
       127060 TCCTTCTCTGGGGCTTATTTTTGTTCAGAACTAGACCAGAGTGTTTGAACCTCCTTTGCA 127001
Db
    Qу
       Db
   Qу
       Db
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Qу
       126880 GGTAATGTGGCATTACTGGCCCACAGAGGTTTTGAGCCAATCAGCTCTGAGACTGGGTTA 126821
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    2641 GAATGTAACAGCTTTAACTTGGGATTTAAGAAGCTTTTAAAAGGTAATAATCCTCTGAAA 2700
Qy
       126820 GAATGTAACAGCTTTAACTTGGGATTTAAGAAGCTTTTAAAAGGTAATAATCCTCTGAAA 126761
Db
Qу
    2701 GAAAAATGACGTAACCACAGCGTGTACTATGAAAGCTGTTATTTTAATAAAGAACGCTGG 2760
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Db
    Qу
       Db
    2881 TCTTTTTTGAGGCAGG 2896
Qу
       11111111111
Db
   126580 TCTTTTTTGAGACAGG 126565
```

Claim 8 is rejected under 35 U.S.C. 103(a) as being unpatentable over GenBank[®] Accession No. AC008736 (September 27, 2000) in view of applicants' admitted state of the prior art (*e.g.*, page 82, first full paragraph). GenBank[®] Accession No. AC008736 has 92.7% sequence identity to SEQ ID NO: 102. Applicants acknowledge the expression of nucleic acids in heterologous host cells to be old (*e.g.*, instant application at page 82, first full paragraph). It would have been obvious for one of ordinary skill in the

art at the time the invention was made to express the nucleic acid of GenBank[®] Accession No. AC008736 in the admittedly old manner in order to produce large amounts of sequence-specific polypeptide. This rejection is repeated for reasons already of record (*e.g.*, Office action mailed |June 2, 2006, page 10). The discussion in the rejection under 35 U.S.C. § 102(b) is incorporated here.

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THIS ACTION IS MADE FINAL. Applicant is reminded of the extension of time policy as set forth in 37 CFR 1.136(a).

A shortened statutory period for reply to this final action is set to expire THREE MONTHS from the mailing date of this action. In the event a first reply is filed within TWO MONTHS of the mailing date of this final action and the advisory action is not mailed until after the end of the THREE-MONTH shortened statutory period, then the shortened statutory period will expire on the date the advisory action is mailed, and any extension fee pursuant to 37 CFR 1.136(a) will be calculated from the mailing date of the advisory action. In no event, however, will the statutory period for reply expire later than SIX MONTHS from the mailing date of this final action.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to James Martinell whose telephone number is (571) 272-0719.

The examiner works a flexible schedule and can be reached by phone and voice mail.

Alternatively, a request for a return telephone call may be e-mailed to james.martinell@uspto.gov. Since e-mail communications may not be secure, it is suggested that information in such requests be limited to name, phone number, and the best time to return the call.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Ram Shukla, can be reached on (571) 272-0735.

OFFICIAL FAX NUMBER

The fax phone number for the organization where this application or proceeding is assigned is (571) 273-8300. Any Official Communication to the USPTO should be faxed to this number.

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Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to (571) 272-0547.

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For all other customer support, please call the USPTO Call Center (UCC) at 800-786-9199.

James Martinell, Ph.D. Primary Examiner

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